=> d his (FILE 'HOME' ENTERED AT 18:34:18 ON 03 JAN 2002) FILE 'MEDLINE, CAPLUS, BIOSIS, SCISEARCH' ENTERED AT 18:34:33 ON 03 JAN 2002 90 S TOTAL (3A) YEAST (W) (RIBONUCLEIC (W) ACID OR RNA) L1327786 S INFLAMMATION L20 S L1 AND L2 L342 DUP REM L1 (48 DUPLICATES REMOVED) L4L5 66 S TOTAL(W) YEAST(W) (RIBONUCLEIC(W) ACID OR RNA) 27 DUP REM L5 (39 DUPLICATES REMOVED) L6 => d au ti so 1-27 16 ANSWER 1 OF 27 BIOSIS COPYRIGHT 2002 BIOSIS L6 ΑU Yarmoluk, S. M. (1); Kovalska, V. B.; Kryvorotenko, D. V.; Balanda, A. 0.; Ogul'chansky, T. Yu. ΤI Interaction of cyanine dyes with nucleic acids. XXV. Influence of affinity-modifying groups in the structure of benzothiazol-4-(2,6dimethylpyridinium) dyes on the spectral properties of the dyes in the presence of nucleic acids. Spectrochimica Acta Part A Molecular and Biomolecular Spectroscopy, SO (June, 2001) Vol. 57A, No. 7, pp. 1533-1540. print. ISSN: 1386-1425. 1.6 ANSWER 2 OF 27 SCISEARCH COPYRIGHT 2002 ISI (R) Zhao H; Bojanowski K; Ingber D E; Panigrahy D; Pepper M S; Montesano R; ΑU Shing Y (Reprint) New role for tRNA and its fragment purified from human urinary bladder ΤI carcinoma conditioned medium: Inhibition of endothelial cell growth JOURNAL OF CELLULAR BIOCHEMISTRY, (JAN 2000) Vol. 76, No. 1, pp. 109-117. Publisher: WILEY-LISS, DIV JOHN WILEY & SONS INC, 605 THIRD AVE, NEW YORK, NY 10158-0012. ISSN: 0730-2312. ANSWER 3 OF 27 CAPLUS COPYRIGHT 2002 ACS 1.6 Kang, John J.; Watson, Robert M.; Fisher, Mary E.; Higuchi, Russell; ΑU Gelfand, David H.; Holland, Michael J. Transcript quantitation in total yeast cellular RNA using kinetic PCR TI Nucleic Acids Res. (2000), 28(2), e2, ii-viii SO CODEN: NARHAD; ISSN: 0305-1048 L6 ANSWER 4 OF 27 DUPLICATE 1 ΑU Trachtulec Z; Forejt J TI Transcription and RNA processing of mammalian genes in Saccharomyces cerevisiae. SO NUCLEIC ACIDS RESEARCH, (1999 Jan 15) 27 (2) 526-31. Journal code: O8L; 0411011. ISSN: 0305-1048. DUPLICATE 2 L6ANSWER 5 OF 27 MEDLINE Zhao H; Bojanowski K; Ingber D E; Panigrahy D; Pepper M S; Montesano R; ΑU Shing Y

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genes of three tested were transcribed both from their sense and

antisense

strands and all tested microsatellite, inter-repetitive and anonymous mouse loci were detected in the YAC clone RNA. An RNA transcript from a well defined intergenic region of two head-to-head oriented mouse genes was detected by RT-PCR and by RNase protection assay. These results indicate the presence of multiple yeast-specific transcription sites in the mouse DNA. 3' RACE experiments demonstrated the inability of the

yeast

to use the mouse polyadenylation signals. Thus, a method for isolation of mammalian exons based on a YAC clone RNA is likely to produce a high background, because the enrichment with mammalian exons in the YAC RNA is low. Nevertheless, YAC clones can serve as in vivo test tubes to study

the

conservation of RNA processing sequences.

L6 ANSWER 7 OF 27 MEDLINE

DUPLICATE 4

AB We quantitate the absolute levels of individual mRNAs per yeast cell by hybridizing total yeast RNA with an excess

of gene-specific 32P-oligonucleotides, and digesting the resulting ${\tt RNA-DNA}$

hybrids with S1 nuclease. By comparing the his3 hybridization signal from a known amount of yeast cells to the signal generated by a known amount of

his3 RNA synthesized in vitro, we determine that yeast strain KY114 growing in yeast extract/peptone/glucose medium at 30 degrees C contains seven molecules of his3 mRNA per cell. Using a galactose shut-off procedure, we determined that the half-life of his3 mRNA is approximately 11 min under these conditions. From these observations, we calculate that one his3 mRNA molecule is synthesized every 140 s. Analysis of other his3 promoter derivatives suggests that the maximal transcriptional initiation rate in yeast cells is one mRNA molecule every 6-8 s. Using his3 as an internal standard, the number of mRNA molecules per cell have been determined for ded1, trp3, rps4, and gall under a variety of growth conditions. From these results, the absolute mRNA level of any yeast gene can be determined in a single hybridization experiment. Moreover, the

rate

of transcriptional initiation can be determined for mRNAs whose decay rates are known.

L6 ANSWER 10 OF 27 MEDLINE

DUPLICATE 7

AB The TIF3 gene of Saccharomyces cerevisiae was cloned and sequenced. The deduced amino acid sequence shows 26% identity with the sequence of mammalian translation initiation factor eIF-4B. The TIF3 gene is not essential for growth; however, its disruption results in a slow growth and

cold-sensitive phenotype. In vitro translation of **total**yeast RNA in an extract from a TIF3 gene-disrupted
 strain is reduced compared with a wild-type extract. The translational
 defect is more pronounced at lower temperatures and can be corrected by
 the addition of wild-type extract or mammalian eIF-4B, but not by
addition

of mutant extract. In vivo translation of beta-galactosidase reporter mRNA

with varying degree of RNA secondary structure in the 5' leader region in a TIF3 gene-disrupted strain shows preferential inhibition of translation of mRNA with more stable secondary structure. This indicates that Tif3 protein is an RNA helicase or contributes to RNA helicase activity in vivo.